



Vaccine

journal homepage: www.elsevier.com/locate/vaccineProtective immunity against tick infestation in cattle vaccinated with recombinant trypsin inhibitor of *Rhipicephalus microplus*Renato Andreotti^a, Rodrigo Casquero Cunha^a, Mariana Aparecida Soares^a, Felix D. Guerrero^b, Fábio P. Leivas Leite^c, Adalberto A. Pérez de León^{b,*}^a EMBRAPA Beef Cattle, Avenida Rádio Maia, 830 – Vila Popular – Caixa Postal 154, CEP 79106-550 – Campo Grande, MS, Brazil^b USDA-ARS Knippling-Bushland U.S. Livestock Insects Research Laboratory, 2700 Fredericksburg Road, Kerrville, TX 78028, USA^c Universidade Federal de Pelotas (UFPEL), Institute of Biology, Department of Microbiology and Parasitology, Rua Gomes Carneiro, 1 – Centro – Caixa Postal 354, CEP 96010-610 – Pelotas, RS, Brazil

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ABSTRACT

The cattle tick, *Rhipicephalus microplus*, is regarded as the most economically important ectoparasite of livestock globally. Control is achieved primarily through the use of acaricides. This approach is hampered by the development of resistance to commercial acaricides among cattle tick populations. Vaccination against *R. microplus* infestation is another technology that can be integrated for effective cattle tick control. Proteins belonging to the Kunitz-BPTI family are abundant in cattle tick salivary glands, midgut, and ovaries. These organs are attractive targets for the development of a novel cattle tick vaccine. Efficacy assessment against cattle tick infestation in bovines using a vaccine containing the recombinant form of a member of the Kunitz family from *R. microplus* produced in a yeast expression system is reported for the first time here. The yeast *Pichia pastoris* was bioengineered to produce the recombinant version of a trypsin inhibitor that is expressed in cattle tick larvae (rRmLTI). Immunization with rRmLTI afforded 32% efficacy against *R. microplus*. The estimated molecular weight of rRmLTI was 46 kDa. Structural homology to the native form of the larval trypsin inhibitor was documented by recognition of rRmLTI in Western-blots using polyclonal antibodies from mice immunized with cattle tick larval extract or rRmLTI. Bioinformatics analysis of the partial nucleotide and deduced amino acid sequences indicated that the rRmLTI closely resembles BmTI-6, which is a three-headed Kunitz protein present in cattle tick ovary and fat tissue.

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1. Introduction

Ectoparasitism of cattle by the southern cattle tick, *Rhipicephalus microplus*, inflicts severe economic losses to the livestock industry. Cattle productivity is undermined by the direct effects of ectoparasitism and indirectly by the role *R. microplus* plays as vector of the infectious agents causing bovine babesiosis and anaplasmosis [1,2]. The control of *R. microplus* is achieved mainly through the use of chemical acaricides [3]. However, chemical acaricides have not been utilized judiciously. This has led to the development of acaricide resistance among populations of *R. microplus* [4,5].

Vaccinating cattle with tick molecules formulated as antigens to elicit a protective immune response is a strategy proven useful for

the integrated control of cattle ticks [7,10,36]. The benefits of using anti-tick vaccines as part of an integrated control program include a reduction in the use of acaricides, extending the useful life of acaricides by delaying the onset of resistance, reducing the incidence of *R. microplus*-borne diseases, and decreased production costs [6,8,9]. The only tick molecule currently developed and marketed as a component of an anti-tick vaccine is Bm86 from *R. microplus*. Bm86 is a glycoprotein expressed in eggs a few days after oviposition, unfed and blood-fed larvae, nymphs, adult males, and in the ovaries of partially engorged adult females [11]. The Bm86 gene appeared to be down-regulated in the ovaries of ticks feeding on cattle infected with *B. bovis* [12]. Anti-tick vaccine products based on the recombinant version of Bm86 (rBm86) were registered in Australia under the trade name TickGARD®, and in Cuba as Gavac® in the 1990s [13,36]. The rBm86-based vaccines are highly efficacious against *R. annulatus* [38,39], but the level of efficacy against *R. microplus* varies according to characteristics of the tick population targeted and host factors among other things [14,15]. Pen trials conducted in the state of Mato Grosso do Sul, Brazil revealed that the efficacy of Bm86-based vaccines against the Campo Grande strain of *R. microplus* ranged from 31 to 49% [17,18].

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Efficacy around 99% against *R. annulatus* obtained with Bm86-based vaccines is an indication of the consistent high level of anti-*R. microplus* immunoprotection that a novel antigenic and immunogenic tick molecule, or combinations thereof, could elicit in vaccinated cattle. Such level of efficacy offers the opportunity to incorporate vaccination as a tool for the integrated eradication of cattle fever tick populations [40,41]. The search for protective antigens that are highly efficacious against *R. microplus* continues. Proteinase inhibitors have received attention as a group of molecules found in ticks with potential for use as immunogens in an anti-tick vaccine. Several trypsin inhibitors that are present in the egg, larval and adult stages of *R. microplus* have been described [19–21]. It has been suggested that the *R. microplus* serine protease inhibitors may be involved in larval attachment at the bite site and blood feeding [22]. Trypsin inhibitors from *R. microplus* larvae purified in their native form elicited a protective immune response in vaccinated cattle yielding 72.8% efficacy, and 69.7% reduction in the number of adult female ticks completing the parasitic phase of their life cycle [22]. However, a peptide designed from one of the *R. microplus* larval trypsin inhibitors afforded only 18.4% immunoprotection against tick infestation in crossbred cattle [23].

The use of recombinant trypsin inhibitors can circumvent the challenge of having to purify trypsin inhibitors in sufficient quantities to conduct cattle tick vaccination tests [21,22]. An expressed sequence tag originally identified in *R. microplus* larvae was later reported to correspond to sequence amplified from ovarian tissue coding for the fragment of a Kunitz-BPTI domain protease inhibitor termed rBmTI-6 [21,24]. The rBmTI-6 was expressed in the *Pichia pastoris* system and characterized as a three-headed Kunitz-bovine pancreatic trypsin inhibitor, but its ability to protect immunized cattle against tick infestation remained to be determined [21]. Here, the partial nucleotide sequence of the putative *R. microplus* larval trypsin inhibitor was used to produce the recombinant polypeptide in the yeast expression system to probe its immunoprotective properties [24]. Results of the cattle immunization trial and other experiments using the recombinant *R. microplus* larval trypsin inhibitor (rRmLTI) are also reported.

2. Materials and methods

2.1. Ticks

Ticks used for this study were obtained from a laboratory colony maintained at EMBRAPA Beef Cattle. This colony was established using ticks collected from bovines in ranches around Campo Grande, Matto Grosso do Sul, Brazil. Ticks were maintained under laboratory conditions for two years prior to use in the experiments reported here. Cattle were used to cycle the tick progeny. Tick stages requiring incubation were kept in the laboratory at 28 °C and 80% relative humidity. The Campo Grande cattle tick strain is susceptible to commercially available acaricides.

2.2. Transformation of *Pichia pastoris* with RmLTI DNA construct and sequence analysis

The expressed sequence tag (EST) coding for RmLTI (GenBank ID: CK186726 [21,24]) was optimized for *P. pastoris* codon usage, and synthesized by Epoch Biolabs, Inc. Codon optimization was done using Epoch Biolabs, Inc. proprietary software set at 15% cut off for codon efficiency. This RmLTI DNA fragment was cloned into pPICZαA, producing the pPICZαRmLTI construct. The recombinant plasmid codes for a His tag that is added to the N-terminus of the protein product. Previously described procedures were followed to produce rRmLTI in the *P. pastoris* expression system [25].

Alignment, similarity, and discordance comparisons based on bioinformatics techniques were conducted between predicted amino acid sequences for: rRmLTI, EST CK186726, BmTI-6 from ovarian cDNA (GenBank ID: P83606.2), and N-terminal amino acid sequence information for BmTI-A (GenBank ID: P83609), BmTI-D (GenBank ID: P83607), BmTI-2 (GenBank ID: P83603), and BmTI-3 (GenBank ID: P83604). ClustalW from the BioEdit suite was used with Vector NTI® software (Invitrogen) as described previously to conduct the bioinformatics analyses [16]. The amino acid sequence from rRmLTI was submitted to protein function and superfamily analysis using the protein domains identifier software InterProScan [42].

2.3. rRmLTI quantitation and purification, and murine antibody production

Protein concentration in *P. pastoris* culture supernatant was quantified as described previously [25]. Proteins were precipitated with methanol and the precipitated proteins resuspended in denaturing binding buffer (8 M Urea, 20 mM sodium phosphate pH 7.8, 500 mM NaCl). The rRmLTI was purified using a Ni²⁺ charged Ni-NTA (Qiagen, Hilden, Germany) affinity column with denaturing elution buffer (8 M Urea, 20 mM Sodium Phosphate pH 4.0, 500 mM NaCl) and the purification process monitored by 7.5% SDS-PAGE. Eluted fractions of high purity were pooled and dialyzed against PBS.

Animal care and use was conducted at EMBRAPA Beef Cattle according to institutional guidelines. Polyclonal serum against *R. microplus* larval extract or rRmLTI was produced using BALB/c mice as described previously [25]. The RmLTI vaccine was prepared with 500 µg of rRmLTI protein resuspended in 4 mL of 150-mM Tris-HCl at pH 7.4 and emulsified with 6 mL of Montanide ISA 61 VG (Seppic, Paris). Twelve female BALB/c mice were used, which were separated into two groups of six animals. One group received the rRmLTI formulation and the other the larval extract preparation. Each mouse within the respective group was immunized with 50 µg mL⁻¹ dose⁻¹ of rRmLTI, or 100 µg mL⁻¹ dose⁻¹ of larval extract. Three subcutaneous doses were applied at 21-day intervals. Sera were collected 21 days after the last dose.

2.4. rRmLTI electrophoresis and Western blotting

Electrophoresis analysis was performed on material from a 10% polyacrylamide gel run with 10 µL of culture supernatant per well containing 0.55 µg µL⁻¹ of protein. Subsequent Western blotting analysis was performed as previously described [25].

2.5. Pen trial, bovine serum analysis, and efficacy assessment

The rRmLTI antigen expressed in *P. pastoris* was adjuvated with Montanide ISA 61 VG (Seppic, Paris) and doses of 2 mL containing 100 µg of the recombinant protein prepared. One-year old Holstein calves were randomly distributed into two groups of six animals each. One group was immunized with rRmLTI antigen purified and formulated as described above. The second group (negative controls) was injected with adjuvant/saline alone. Serum samples were collected and processed, and all procedures involving ticks were performed according to methods described previously [17]. Sera obtained immediately before the initial injection, and at different time points thereafter, from each of the six cattle in the vaccinated and control groups were pooled and stored in an ultralow freezer until ready for ELISA testing.

For ELISA, microtiter plates were coated with 1 µg mL⁻¹ of rRmLTI antigen in 20 mM sodium carbonate buffer (pH 9.6), 50 µL per well, and incubated overnight at 4 °C. Duplicate samples of

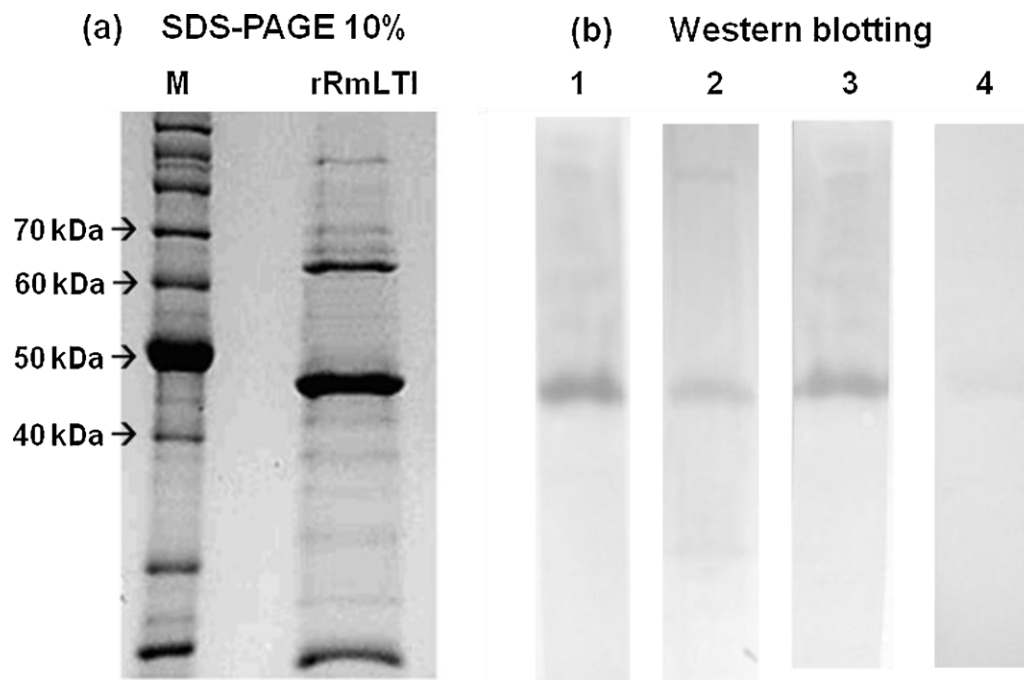


Fig. 1. Identification of *P. pastoris* product as larval trypsin inhibitor present in *R. microplus* larvae by electrophoresis and immunoblotting. Supernatant of culture containing *P. pastoris* transformed with pPICZαA-RmLTI was processed to perform: (a) electrophoresis (SDS-PAGE 10%) where the gel was loaded with M (molecular weight marker [BenchMark™ Protein Ladder, Invitrogen]), and rRmLTI (supernatant of *P. pastoris* culture expressing rRmLTI); (b) Western blot of rRmLTI detected with: (1) anti-His monoclonal antibody, (2) serum from mice vaccinated with rRmLTI, (3) serum from mice vaccinated with larval extract, and (4) serum from unvaccinated mice.

pooled sera for each group and sampling date were tested. Subsequent procedures were performed as described previously [25].

Procedures described before were followed to assess treatment effects on tick biology and vaccine efficacy [17,26]. Engorged female ticks dropping off of cattle from each group were collected daily for 13 days and incubated in the laboratory to obtain eggs that were pooled until 1 gram was accumulated. The egg masses obtained per collection day from several ticks in each group were incubated to determine hatching rates.

2.6. Ex vivo feeding of anti-rRmLTI IgG and assessment of tick egg eclosion rates

Serum samples from bovines immunized with rRmLTI were collected just prior to tick infestation and subjected to affinity chromatography on a protein A-Sepharose column to purify immunoglobulin G (IgG). The IgG eluted from the column resulted in a yield of 5 mg mL⁻¹ of non-immune serum. Four groups, each consisting of ten adult female ticks, were set up for treatment. Each tick in the respective group was fed with the antibody mixture containing 0, 25, 50, or 100 µg of purified IgG in 20 µL by placing a capillary tube in its hypostome. The effect on egg eclosion was assessed as described previously [17].

2.7. Statistics

Data on female reproductive parameters were analyzed using a *t*-test. Otherwise, differences between means were determined using one-way analysis of variance (ANOVA). Differences were considered significant when *p* < 0.05.

3. Results

3.1. Production of rRmLTI in *P. pastoris* expression system

Identity of the DNA insert in pPICZαRmLTI was confirmed by sequencing and alignment with the RmLTI clone sequence. One

Mut⁺ clone was selected and analysis of the induced recombinant protein revealed a band of approximately 46 kDa. The calculated molecular weight for rRmLTI was 37.9 kDa. It remains to be validated if post-translational modification through glycosylation, potentially adding 8–9 mannose and two N-acetyl-glucosamine molecules, may have caused the discrepancy between the calculated and observed molecular weights of the protein product in *P. pastoris*. Direct quantification from culture supernatants revealed rRmLTI production levels of 550 mg L⁻¹. Analysis of the nickel column purification product showed a protein of 46 kDa and the yield following purification was 870 mg L⁻¹. Western blot analysis of the rRmLTI protein was carried out with primary sera from mice (anti-*R. microplus* larval extract and anti-rRmLTI) and anti-His tag monoclonal antibody revealing affinity for a protein of approximately 46 kDa (Fig. 1).

3.2. Immunoprotection against *R. microplus* infestation in cattle vaccinated with rRmLTI

The antibody response of cattle immunized with the vaccine formulation containing rRmLTI is shown in Fig. 2. Antibody levels against rRmLTI peaked around 31 days after the second booster immunization. Tick infestations were established around ten days before the apparent decline in the specific antibody response commenced. A transient effect on the average weight of engorged adult female ticks dropping off of vaccinated cattle was apparent through the ninth day of the collection period (Fig. 3). With the exception of days 2 and 4, the average weight of engorged female ticks collected from the vaccinated group was significantly lower up to day nine (Fig. 3; *p* < 0.05). Equivalence of the average engorged adult female tick weight between groups beyond day 9 of the collection period was temporally associated with the aforementioned decline in anti-rRmLTI antibody levels (Fig. 2). A similar tendency was observed in the eclosion rate for eggs collected from ticks detaching from vaccinated cattle (Fig. 4).

Table 1Efficacy of vaccine containing recombinant *R. microplus* larval trypsin inhibitor against ticks infesting cattle and its effect on female reproductive parameters.

Animal	Tick total number		Tick mean weight (mg)		Egg mean weight (mg)		Larval hatchability (%)	
	Control	Vaccinated	Control	Vaccinated	Control	Vaccinated	Control	Vaccinated
1	1209	818	267	210	112	161	95	90
2	661	441	254	209	138	142	98	85
3	571	588	276	209	121	143	97	86
4	521	602	285	234	145	160	98	89
5	1089	671	304	209	160	154	96	91
6	543	89	275	190	149	125	96	87
Mean ± SD ^a	766 ± 303	535 ± 250	277 ± 17	210 ± 14	138 ± 18	148 ± 014	97 ± 1.2	88 ± 2.4
<i>t</i> -Test	<i>p</i> = 0.1810		<i>p</i> < 0.0001		<i>p</i> = 0.3040		<i>p</i> < 0.0001	
% Reduction ^b	DT = 30.15		DW = 24.66		DO = −7.27		DF = 8.97	
Efficacy = 100 [1 − (535/766 × 148/138 × 88/97)] = 32%								

^a Arithmetic mean \pm standard deviation; *p*-values of *t*-test for independent samples are shown.^b Percent reduction was calculated in relation to the control unvaccinated group: DT, adult female ticks; DW, tick weight; DO, egg laying capacity; DF, fertility. Efficacy (%) = 100 [1 - (CRT \times CRO \times CRF)]; where CRT: reduction in the number of adult female ticks, CRO: reduction in the egg laying capacity, CRF: reduction in fertility.

The cumulative count of engorged adult female ticks collected up to day 13 after detachment started was used to calculate the effects of vaccination with rRmLTI (Table 1). Vaccinated cattle had 30% less ticks detaching from them than the animals injected with adjuvant only. Although egg laying capacity was unaffected, there was a significant effect associated with vaccination on tick weight and larval hatchability (Table 1; $p < 0.05$). Overall, the rRmLTI vaccine afforded 32% immunoprotection against cattle tick infestation (Table 1).

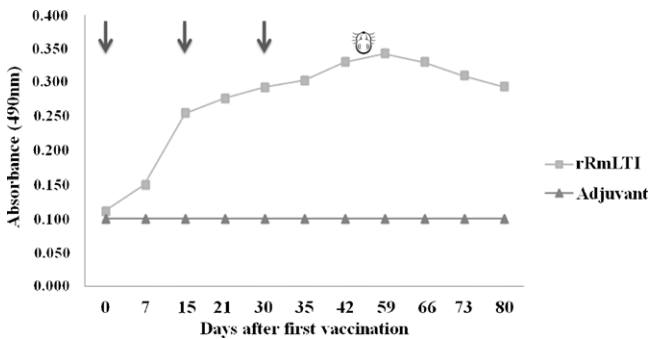


Fig. 2. Profile of humoral immune response in cattle vaccinated with rRmLTI. Absorbance readings are for ELISA tests of weekly antisera samples collected from two groups of six cattle immunized at 0, 15, and 30 days (arrows) with rRmLTI (squared line) or adjuvant (triangled line). Tick icon indicates day 51 after the first vaccination, which is when cattle were infested. Antibody titers of immunized cattle are depicted as the OD_{490 nm} value of the 1:100 dilution of serum samples. Individual samples in each group collected on the day shown in the figure were pooled for ELISA testing.

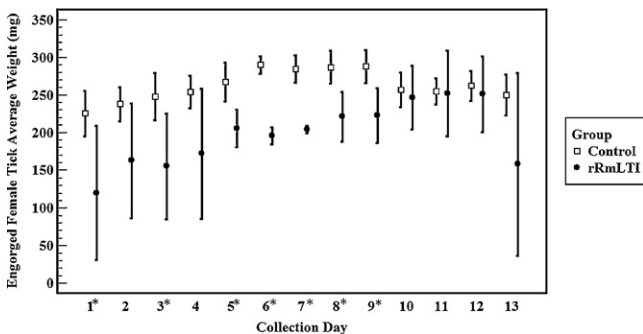


Fig. 3. Reduced weight of female *R. microplus* ticks feeding on cattle vaccinated with rRmLTI. *Group means \pm SD are shown. Collection day with asterisk indicates the average weight between engorged female ticks feeding on vaccinated or control cattle was significantly different ($p < 0.05$).

3.3. Effect of purified anti-rRmLTI IgG on tick egg eclosion

The effect of the anti-rRmLTI antibody response on egg hatching was explored further *ex vivo*. An inverse dose-response was observed between egg hatching and the amount of IgG imbibed by the gravid tick (Fig. 5). The viability of eggs laid by female ticks ingesting IgG antibodies from cattle vaccinated with rRmLTI was significantly compromised and hatching decreased 75.6% in eggs from ticks fed 100 μ g of IgG ($p < 0.05$).

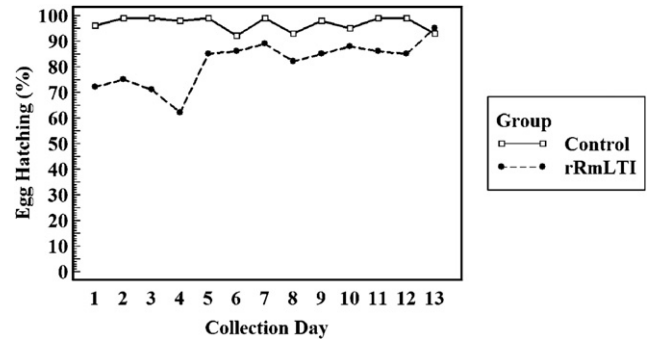


Fig. 4. Percent hatch of eggs laid by female *R. microplus* ticks feeding on control cattle or those vaccinated with rRmLTI. Refer to text for details on egg processing.

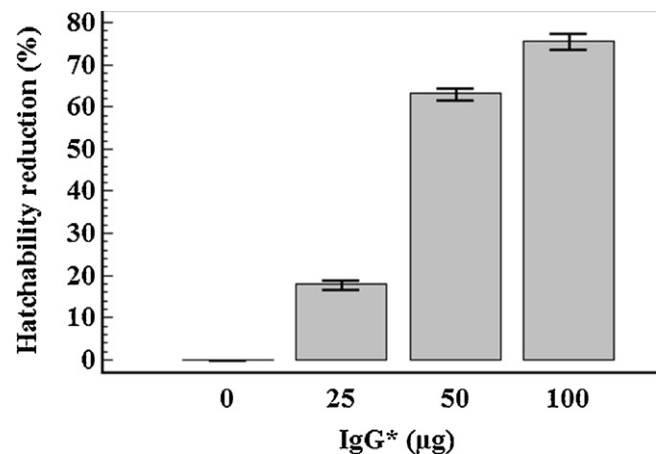


Fig. 5. Anti-rRmLTI IgG imbibed by female cattle ticks inhibits egg hatching in a dose-dependent manner. *Four groups of ten ticks received 0, 25, 50 and 100 μ g of purified IgG. Each tick was fed the specified amount of IgG in a volume of 20 μ L using a capillary tube. Treatment means \pm SD are shown. Mean percent hatch inhibition was significantly different between the treatment groups.

rRmLTI	EAEAEFFETYCKPTYDSGPCKGYFPRNWFNVKTGQCEEFIYGGCQGNKNNHVTRKECETR	60
BmTI-6	----VDFETYCKPTYDSGPCKGYFPRNWFNVKTGQCEEFIYGGCQGNKNNHVTRKECETR	56
CK186726.1	----VDFETYCKPTYDSGPCKGYFPRNWFNVKTGQCEEFIYGGCQGNKNNHVTRKECETR	56
BmTI-2	-----	
BmTI-A	-----	
BmTI-3	-----	
BmTI-D	-----	
rRmLTI	CLRKQLSRLHFSPPVDQYPYGDTERSNEEVPEYPPVHLNDSLEVSAMNQRPLRNYTKQH	120
BmTI-6	CLRKQLSRLHFSPPVDQYPYGDTERSNEEVPEYPPVHLNDSLEVSAMNQRPLRNYTKQH	116
CK186726.1	CLRKQLSRLHFSPPVDQYPYGDTERSNEEVPEYPPVHLNDSLEVSAMNQRPLRNYTKQH	116
BmTI-2	-----	
BmTI-A	-----SQPHVNPFA-----CYVAPDQGPCRAIL---	23
BmTI-3	-----	
BmTI-D	-----	
rRmLTI	KPNVTSDFSASISLTSGVDFETYCKPTHDRGPCKAYIPRNWFNVKTGQCEQFIYGGCQGNK	180
BmTI-6	KPNVTSDFSASISLTSGVDFETYCKPTHDRGPCKAYIPRNWFNVKTGQCEQFIYGGCQGNK	176
CK186726.1	KPNVTSDFSASISLTSGVDFETYCKPTHDRGPCKAYIPRNWFNVKTGQCEQFIYGGCQGNK	176
BmTI-2	--VTIGPV-----CELPKEVGGPCRGHIIIPRY-----	25
BmTI-A	--RYFDDD-----TQTCQRFYGGCEGNANNXXXXEQC-----KAS	58
BmTI-3	-EVHNF-----CLGKPDPGGC--AHYIYRRY-----Y	25
BmTI-D	--AVDFSQV?-----TATAGPCKG-----	17
rRmLTI	NNYETKSICETNCLRRQLSELGVSADVHYRKHNNETKYSNVTVEYPAVHFNVTLNPFVCN	240
BmTI-6	NNYETKSICETNCLRRQLSELGVSADVHYRKHNNETKYSNVTVEYPAVHFNVTLNPFVCN	236
CK186726.1	NNYETKSICETNCLRRQLSELGVSADVHYRKHNNETKYSNVTVEYPAVHFNVTLNPFVCN	236
BmTI-2	-----	
BmTI-A	CKPETEY-----EAKKCLARPES-----GPCLAYMPMNG-----	87
BmTI-3	YV-----	27
BmTI-D	-----	
rRmLTI	EPKYPCLCKGYFPRYYYNSRSKTCCKFIYGGCQSNNGNFLTLEECENTCLVDLQVPRAA	300
BmTI-6	EPKYPCLCKGYFPRYYYNSRSKTCCKFIYGGCQSNNGNFLTLEECENTCLVDLQV----	291
CK186726.1	EPKYPCLCKGYFPRYYYNSRSKTCCKFIYGGCQSNNGNFLTLEECENTCLVDLQ----	290
BmTI-2	-----	
BmTI-A	-----YDSKLGQCVEFIYGGCDGNDNKYTTEEECLKSCK-----	121
BmTI-3	-----	
BmTI-D	-----	
rRmLTI	AAASFLEQKLISEEDLNSAVDHHHHHH	327
BmTI-6	-----	
CK186726.1	-----	
BmTI-2	-----	
BmTI-A	-----	
BmTI-3	-----	
BmTI-D	-----	

Fig. 6. Alignment of predicted amino acid sequences for rRmLTI, EST CK186726, BmTI-6 from ovarian cDNA, and N-terminal amino acid sequence information for BmTI-A, BmTI-D, BmTI-2, and BmTI-3.

3.4. Identification of RmLTI as a member of the Kunitz-bovine pancreatic trypsin inhibitor family

A comparison of the DNA sequences from the EST CK186726 and the RmLTI clone optimized for codon usage in *P. pastoris* revealed 77% identity between the two sequences. The RmLTI DNA sequence in the yeast expression system was missing nineteen bases of the corresponding EST sequence (data not shown).

Fig. 6 depicts the alignment of predicted amino acid sequences for rRmLTI, EST CK186726, BmTI-6 from ovarian cDNA, and N-terminal amino acid sequence information for BmTI-A, BmTI-D, BmTI-2, and BmTI-3. The deduced amino acid sequences between rRmLTI, EST CK186726, and BmTI-6 are 99% identical. Nucleic acid sequence coding for six additional amino acids (EAEAEF) in the N-terminus, and thirty two amino acids (VPRAAAAASFLEQKLISEEDLNSAVDHHHHHH) in the C-terminal portion of the putative

rRmLTI product was added during cloning procedures to allow insertion of a restriction site and coding sequence for the poly-His peptide.

The similarity between their partial amino acid sequences suggested that RmLTI in larvae is a member of the Kunitz-bovine pancreatic trypsin inhibitor (BPTI) family like BmTI-6 in the ovary of adult female cattle ticks. Further exploration of the putative function of RmLTI is reflected in Fig. 7. Relevant protein signature features identified in the deduced amino acid sequence encoded in the expressed sequence tag CK186726 include three putative Kunitz-BPTI domains and two putative Kunitz proteinase inhibitor I2 conserved sites. As noted in BmTI-6, six N-glycosylation sites were present in the partial protein sequence of RmLTI. Six cysteine residues were observed within each of the three Kunitz domains, which are thought to form disulfide linkages contributing stability to the compact polypeptide in its folded form.

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1  GAG GCT GAA GCT GAA TTC TTC GAG ACT TAC TGC AAA CCA ACT TAC GAC TCT GGT CCA TGT
1  E  A  E  A  E  F  F  E  T  Y  C  K  P  T  Y  D  S  G  P  C
61  AAA GGT TAC TTC CCT AGA TGG TGG TTT AAT GTT AAG ACT GGT CAG TGC GAA GAG TTC ATC
21  K  G  Y  F  P  R  W  W  F  N  V  K  T  G  Q  C  E  E  F  I
121 TAC GGT GGT TGC CAA GGT AAC AAG AAC AAC CAT GTT ACT AGA AAG GAG TGC GAA ACC AGA
41  Y  G  G  C  Q  G  N  K  N  N  H  V  T  R  K  ↑  E  C  E  T  R
181 TGT TTG AGA AAG CAA TTG TCC AGA TTG CAT TTC TCC CCA CCT GTC GAC CAA TAC CCT TAC
61  C  L  R  K  ↑  Q  L  S  R  L  H  F  S  P  P  V  D  Q  Y  P  Y
241 GGA GAT ACC GAG CGT TCC AAC GAG GAG GTT CCT GAA TAC CCA CCA GTC CAC TTG AAC GAC
81  G  D  T  E  R  S  N  E  E  V  P  E  Y  P  P  V  H  L  ..N*..D
301 TCT TTG GAG GTT TCC GCT GTT ATG AAC CAG AGA CCA TTG AGA AAC TAC ACT AAG CAG CAT
101 ..S  L  E  V  S  A  V  M  N  Q  R  P  L  R  ..N*..Y..T..  K  Q  H
361 AAG CCA AAC GTT ACT TCC GAC TTC TCT GCT ATT TCC TTG ACC TCC GGT GTT GAT TTC GAA
121 K  P  ..N*..V..T..  S  D  F  S  A  I  S  L  T  S  G  V  D  F  E
421 ACT TAC TGC AAG CCA ACC CAC GAC AGA GGT CCA TGC AAG GCT TAC ATC CCA CGT TGG TGG
141 T  Y  C  K  P  T  H  D  R  G  P  C  K  A  Y  I  P  R  W  W
481 TTC AAC GTC AAG ACC GGT CAA TGT GAG CAA TTC ATC TAC GGT GGT TGT CAA GGT AAC AAG
161 F  N  V  K  T  G  Q  C  E  Q  F  I  Y  G  G  C  Q  G  N  K
541 AAC AAC TAC GAG ACC AAG TCT ATC TGT GAA ACT AAC TGT TTG CGT CGT CAA TTG TCC GAG
181 N  N  Y  E  T  K  S  I  C  E  T  N  C  L  R  R  ↑  Q  L  S  E
601 TTG GGT GTT TCT GCC GAC GTT CAT TAC AGA AAG CAC TGG AAC GAG ACT AAG TAC TCT CCA
201 L  G  V  S  A  D  V  H  Y  R  K  ↑  H  W  ..N*..E..T..  K  Y  S  P
661 AAC GTC ACT GTC GAG TAC CCA GCC GTT CAC TTC AAC GTC ACC TTG AAC CCA GTC TGC AAC
221 ..N*..V..T..  V  E  Y  P  A  V  H  F  ..N*..V..T..  L  N  P  V  C  N
721 GAA CCA AAG TAC CCA GAG TTG TGT AAG GGT TAC TTC CCA AGA TAC TAC TAC AAC TCC AGA
241 E  P  K  Y  P  E  L  C  K  G  Y  F  P  R  Y  Y  Y  N  S  R
781 TCT AAG ACC TGT AAG AAG TTC ATC TAC GGT GGA TGT CAG TCT AAC GGA AAC AAC TTC TTG
261 S  K  T  C  K  K  ↑  F  I  Y  G  G  C  Q  S  N  G  N  N  F  L
841 ACC TTG GAA GAA TGT GAA AAC ACT TGT TTG GTC GAC TTG CAA GTG GTA CCT CGA GCC GCG
281 T  L  E  E  C  E  N  T  C  L  V  D  L  Q  V  V  P  R  A  A
901 GCG GCC GCC AGC TTT CTA GAA CAA AAA CTC ATC TCA GAA GAG GAT CTG AAT AGC GCC GTC
301 A  A  A  S  F  L  E  Q  K  L  I  S  E  E  D  L  N  S  A  V
961 GAC CAT CAT CAT CAT CAT CAT TGA
321 D  H  H  H  H  H  H  Stop

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Fig. 7. A trypsin inhibitor in *R. microplus* larvae is encoded in expressed sequence tag CK186726. The amino acid sequence deduced from the RmLTI DNA clone optimized for use in *P. pastoris* was scanned for protein signature features using the InterProScan informatics tool. Asparagine residues (N) predicted to be N-glycosylated are marked with asterisks. A dotted line underlines Asn-Xaa-Ser/Thr codons. Cysteine residues (C) predicted to form disulfide bonds are highlighted in gray. Amino acids encompassing putative Kunitz-BPTI domains are underlined. Two predicted Kunitz proteinase inhibitor I2 conserved sites are double-underlined.

4. Discussion

Assessment of the efficacy against cattle tick infestation in bovines using a vaccine containing the recombinant form of a member of the Kunitz-BPTI family from *R. microplus* produced in the *P. pastoris* expression system is reported for the first time here. A specific and robust humoral immune response against rRmLTI was achieved with the vaccination protocol consisting of three immunizations, each applied every two weeks. The 32% efficacy obtained with the rRmLTI formulation reflects the significant challenge of discovering highly efficacious antigens protecting cattle against *R.*

microplus infestation. Vaccination experiments where *Bos indicus* cattle were immunized with a mixture of purified larval trypsin inhibitors containing one or two Kunitz-type domains afforded 72.8% efficacy against *R. microplus* infestation [22]. In contrast, the level of immunoprotection obtained in crossbred cattle vaccinated with the synthetic polypeptide containing 29 aa residues derived from the N terminus of the *R. microplus* trypsin inhibitor A was 18.4% [23]. As the gene encoding RmLTI remains to be fully characterized, the apparent discrepancy between specific antibody levels and the low level of efficacy obtained with the rRmLTI vaccine may be due to the partial gene sequence of the EST used to produce

the recombinant protein product in the yeast expression system. Alternatively, the structural and functional redundancy in proteins belonging to the Kunitz family present in *R. microplus* may also account for the relatively low efficacy attained with the rRmLTI vaccine.

Although binding of rRmLTI by polyclonal antibodies from mice immunized with tick larva extract indicates that the recombinant polypeptide produced in *P. pastoris* was as antigenic as the native form of the cognate larval trypsin inhibitor, it is possible that those antibodies recognized epitopes shared by the several trypsin inhibitors discovered in *R. microplus* larvae. Antiserum from cattle vaccinated with purified *R. microplus* trypsin inhibitors recognized rBmTI-6 produced in *P. pastoris* [21]. Antigenic similarity apparently extends beyond intra-specific boundaries because antiserum against the native form of *R. microplus* larval trypsin inhibitors cross-reacts with trypsin inhibitors identified in *R. sanguineus* larvae [27]. Immunogenicity of the rRmLTI is reflected in the kinetics of the bovine humoral immune response. The significant effect on the rate of larvae hatching from eggs laid by female ticks parasitizing vaccinated cattle, which was amplified by feeding female ticks with purified anti-rRmLTI IgG suggests that potentiation of the humoral response, perhaps using other adjuvants, could enhance the efficacy of a polyvalent vaccine with Kunitz inhibitors from *R. microplus*. Adjuvant choice was shown to influence antibody levels, which correlated with the level of inhibition on malaria parasites [28]. However, no direct correlation was observed between antibodies against rRmLTI and overall efficacy in our study. By comparison, the vast array of Kunitz type inhibitors present in *R. microplus* was invoked to explain the apparently small impact silencing the gene coding for boophilin, a double Kunitz type thrombin inhibitor expressed in the gut, had on egg production [29].

Considering the purported involvement of larval trypsin inhibitors and confirmed role of other Kunitz inhibitors in blood feeding, the reduced number of female ticks detaching from vaccinated cattle may reflect the impact of bovine anti-rRmLTI antibodies on the ability of *R. microplus* to acquire a blood meal [20,29]. However, the physiological roles of RmLTI and BmTI-6 remain to be determined in the larval and adult stages of the cattle tick, respectively, despite similarities in their partial nucleotide and amino acid sequences. Without knowing the function of RmLTI and BmTI-6, it remains possible that the decrease in hatching rates observed in eggs laid by female ticks fed purified IgG antibodies obtained from vaccinated cattle resulted from the effects of antibody binding to epitopes shared by rRmLTI and the native form of BmTI-6 in *R. microplus* ovaries.

The Kunitz family of polypeptides is one of at least 20 families belonging to the canonical type of serine protease inhibitors [30]. A characteristic of proteins belonging to this family is the Kunitz domain that can be present in single or multiple copies. At least 303 Kunitz proteins have been identified in ticks thus far and some of them can contain as many as seven Kunitz domains [31]. Cysteine patterns were used to classify Kunitz proteins into three groups to facilitate study of their evolution in the Lyme disease vector *Ixodes scapularis* and to expand the analysis to ticks in general [31]. Although primarily involved in proteinase inhibition, the Kunitz domain has evolved to perform other functions requiring protein-protein interactions [32]. Cattle tick ovaries, fat body, hemocytes, and midgut contain Kunitz proteins [21,29,33,34]. Proteomic studies revealed the presence of Kunitz proteins that are up-regulated in ovarian tissue when *R. microplus* is infected with *Babesia bovis* [35]. A publicly available genomic database called CattleTickBase offers the opportunity to study the evolutionary history of Kunitz proteins in *R. microplus* [35]. It is possible that BmTI-6 and the RmLTI encoded by CK186726 are splice variants of the same gene or paralogs of the same Kunitz protein as suggested before

for BmTI-A and other Kunitz proteins present in cattle tick ovary [34].

Previous research documenting 72.8% efficacy against *R. microplus* infestation using purified trypsin inhibitors and the critical role Kunitz proteins play in various biological processes including proteinase inhibition warrant continued vaccine discovery research with this protein family. Production of rRmLTI in *P. pastoris* facilitates its use to formulate polyvalent cattle tick vaccines that include other Kunitz proteins or different antigens from *R. microplus*. The level of immunoprotection attained through vaccination with rRmLTI was low as compared to other novel antigens discovered recently [37,43]. Of note are the results from vaccination using immunogenic peptides that yielded tick efficacy between 80 and 90% [44,45]. Salivary glands, midgut, and ovaries are prime targets to disrupt cattle tick biology using vaccines and Kunitz proteins are abundant in those tissues. The use of epitopes from Kunitz proteins in combination with immunogenic portions of other tick molecules to produce a dual action vaccine could be another way to exploit the redundancy of *R. microplus* Kunitz inhibitors to innovate a highly efficacious cattle tick vaccine.

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